RAW SEQUENCE LISTING PATENT APPLICATION US/08/875,849

DATE: 09/15/98 TIME: 15:05:55

INPUT SET: S28646.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

	1	SEQUENCE LISTING	~
	2 3	(1) General Information:	
	4		The second secon
>	5	(i) APPLICANT:	·
>	6	(A) NAME: LeukoSite, Inc.	142
>	7	(B) STREET: 215 First Street	
>	8	(C) CITY: Cambridge	
>	9 10	(D) STATE/PROVINCE: Massachusetts (E) COUNTRY: U.S.A.	
>	11	(F) POSTAL CODE/ZIP: 02142	
>	12	(G) TELEPHONE: (617) 621-9350	
>	13	(I) TELEFAX: (617) 621-9349	
/	14	(I) IBMETAA. (017) 021-9349	
>	15	(i) APPLICANT/INVENTOR:	
>	16	(A) NAME: Michael J. Briskin	
>	/ 17	(B) STREET: 28 Harbell Street	
>_\	(18	(C) CITY: Lexington	
5	19	(D) STATE/PROVINCE: Massachusetts	
_ _ /`	20	(E) COUNTRY: U.S.A.	
>	21	(F) POSTAL CODE/ZIP: 02173	
	22	• •	
>	23	(i) APPLICANT/INVENTOR:	
>	24	(A) NAME: Douglas J. Ringler	
>	25	(B) STREET: 382 Ocean Avenue, #1008	
>	26	(C) CITY: Revere	
>	27	(D) STATE/PROVINCE: Massachusetts	
>	28	(E) COUNTRY: U.S.A.	
>	29	(F) POSTAL CODE/ZIP: 02151	
	30		
>	31	(i) APPLICANT/INVENTOR:	
>	32	(A) NAME: Dominic Picarella	
>	33	(B) STREET: 2 North Bennet Court, #4	
>	34	(C) CITY: Boston	
>	35	(D) STATE/PROVINCE: Massachusetts	
>	36	(E) COUNTRY: U.S.A.	
>	37	(F) POSTAL CODE/ZIP: 02113	
	38	(i) approaum/rangemmon.	
> >	39 40	(i) APPLICANT/INVENTOR:	
>	40 41	(A) NAME: Walter Newman (B) STREET: 3 Durham Street, #3	
>	41	(B) STREET: 3 Durnam Street, #3 (C) CITY: Boston	
>	43	(D) STATE/PROVINCE: Massachusetts	
>	44	(E) COUNTRY: U.S.A.	
>	45	(F) POSTAL CODE/ZIP: 02115	
/	46	(r) robing Come/ air. Uairs	
	40		

RAW SEQUENCE LISTING PATENT APPLICATION US/08/875,849

DATE: 09/15/98 TIME: 15:05:56

4/		
48	(ii)	TITLE OF INVENTION: Mucosal Vascular Addressins and Uses
49		Thereof
50	, , , , ,	WINDER OF GROUPINGES 12
51	(111)	NUMBER OF SEQUENCES: 13
52 53	(i **)	CORRESPONDENCE ADDRESS:
53 54	(10)	(A) ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
55		(B) STREET: Two Militia Drive
56		(C) CITY: Lexington
57		(D) STATE: Massachusetts
58		(E) COUNTRY: U.S.A.
59		(F) ZIP: 02173-4799
60		
61		
62		
63		
64		
65	(V)	COMPUTER READABLE FORM:
66		(A) MEDIUM TYPE: Floppy disk
67		(B) COMPUTER: IBM PC compatible
68 69		<pre>(C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTWARE: PatentIn Release #1.0, Version #1.30</pre>
70		(D) SOFTWARE: Patentin Release #1.0, Version #1.50
71	(vi)	CURRENT APPLICATION DATA:
72	(/	(A) APPLICATION NUMBER:
73		(B) FILING DATE:
74		(C) CLASSIFICATION:
75		
76	(vii)	PRIOR APPLICATION DATA:
77		(A) APPLICATION NUMBER: US 08/523,004
78		(B) FILING DATE: 01-SEP-1995
79		
80	(V11)	PRIOR APPLICATION DATA:
81 82		(A) APPLICATION NUMBER: US 08/386,857 (B) FILING DATE: 10-FEB-1995
83		(B) FILING DATE: IU-FEB-1995
84	(viii)	ATTORNEY/AGENT INFORMATION:
85	(1111)	(A) NAME: Brook, David E.
86		(B) REGISTRATION NUMBER: 22,592
87		(C) REFERENCE/DOCKET NUMBER: LKS94-04A2 PCT
88		
89	(ix)	TELECOMMUNICATION INFORMATION:
90		(A) TELEPHONE: 617-861-6240
91		(B) TELEFAX: 617-861-9540
92		
93		
94	/0\ TNEO:	DANGEON GOD GEO TO NO. 1.
95 96	(Z) INFO	RMATION FOR SEQ ID NO:1:
96 97	/ i \	SEQUENCE CHARACTERISTICS:
98	(+)	(A) LENGTH: 1624 base pairs
99		(B) TYPE: nucleic acid

RAW SEQUENCE LISTING PATENT APPLICATION US/08/875,849

DATE: 09/15/98 TIME: 15:05:57

								_	_					IN	<i>NPUT</i>	SET:	S28646.raw
100			•	•	TRAN				ole								
101			(]	D) T	OPOL	OGY:	line	ear									
102																	
103		(ii) MO	LECU	LE T	YPE:	CDN	A									
104																	
105																	
106	(ix) FEATURE:																
107			•		AME/I												
108			(]	B) L(OCAT	ION:	1	1218									
109																	
110																	
111																	
112		(xi) SE	QUEN	CE DI	ESCR	IPTI	ON:	SEQ :	ID NO	0:1:						
113																	
114	ATG	GAT	TTC	GGA	CTG	GCC	CTC	CTG	CTG	GCG	GGG	CTT	CTG	GGG	CTC	CTC	48
115	Met	Asp	Phe	Gly	Leu	Ala	Leu	Leu	Leu	Ala	Gly	Leu	Leu	Gly	Leu	Leu	
116	1				5					10					15		
117																	
118	CTC	GGC	CAG	TCC	CTC	CAG	GTG	AAG	CCC	CTG	CAG	GTG	GAG	CCC	CCG	GAG	96
119	Leu	Gly	Gln	Ser	Leu	Gln	Val	Lys	Pro	Leu	Gln	Val	Glu	Pro	Pro	Glu	
120				20					25					30			
121																	
122	CCG	GTG	GTG	GCC	GTG	GCC	TTG	GGC	GCC	TCG	CGC	CAG	CTC	ACC	TGC	CGC	144
123	Pro	Val	Val	Ala	Val	Ala	Leu	Gly	Ala	Ser	Arg	Gln	Leu	Thr	Cys	Arg	
124			35					40					45				
125																	
126	CTG	GCC	TGC	GCG	GAC	CGC	GGG	GCC	TCG	GTG	CAG	TGG	CGG	GGC	CTG	GAC	192
127	Leu	Ala	Cys	Ala	Asp	Arg	Gly	Ala	Ser	Val	Gln	Trp	Arg	Gly	Leu	Asp	
128		50					55					60					
129																	
130	ACC	AGC	CTG	GGC	GCG	GTG	CAG	TCG	GAC	ACG	GGC	CGC	AGC	GTC	CTC	ACC	240
131	Thr	Ser	Leu	Gly	Ala	Val	Gln	Ser	Asp	Thr	Gly	Arg	Ser	Val	Leu	Thr	
132	65					70					75					80	
133																	
134	GTG	CGC	AAC	GCC	TCG	CTG	TCG	GCG	GCC	GGG	ACC	CGC	GTG	TGC	GTG	GGC	288
135	Val	Arg	Asn	Ala	Ser	Leu	Ser	Ala	Ala	Gly	Thr	Arg	Val	Cys	Val	Gly	
136					85					90					95		
137																	
138					CGC												336
139	Ser	Cys	Gly	_	Arg	Thr	Phe	Gln		Thr	Val	Gln	Leu		Val	Tyr	
140				100					105					110			
141																	
142					CAG												384
143	Ala	Phe		Asp	Gln	Leu	Thr		Ser	Pro	Ala	Ala		Val	Pro	Gly	
144			115					120					125				
145																	
146					GCC												432
147	Asp		Glu	Val	Ala	Cys		Ala	His	Lys	Val		Pro	Val	Asp	Pro	
148		130					135					140					
149																	
150					TTC												480
151		Ala	Leu	Ser	Phe		Leu	Leu	Val	Gly	-	Gln	Glu	Leu	Glu	-	
152	145					150					155					160	

RAW SEQUENCE LISTING PATENT APPLICATION US/08/875,849

DATE: 09/15/98 TIME: 15:05:58

														1 P	VPUI	3E1: 3	28040.raw
153 154	GCG	CAA	GCC	CTG	GGC	CCG	GAG	GTG	CAG	GAG	GAG	GAG	GAG	GAG	CCC	CAG	528
155	Ala	Gln	Ala	Leu	Gly	Pro	Glu	Val	Gln	Glu	Glu	Glu	Glu	Glu	Pro	Gln	
156					165					170					175		
157	~~~																
158							TTC										576
159 160	GIY	ASP	GIU	180	vaı	Leu	Phe	arg	185	THE	GIU	arg	тгр	190	Leu	Pro	
161				100					105					190			
162	CCC	CTG	GGG	ACC	ССТ	GTC	CCG	CCC	GCC	СТС	TAC	TGC	CAG	GCC	ACG	ATG	624
163							Pro										
164			195					200			3	2	205				
165																	
166	AGG	CTG	CCT	GGC	TTG	GAG	CTC	AGC	CAC	CGC	CAG	GCC	ATC	CCC	GTC	CTG	672
167	Arg		Pro	Gly	Leu	Glu	Leu	Ser	His	Arg	Gln	Ala	Ile	Pro	Val	Leu	
168		210					215					220					
169	~.~						-1-							~~~	~-~	~~~	=00
170							GAG										720
171 172	225	Ser	Pro	Thr	ser	230	Glu	Pro	Pro	ASP	235	Thr	Ser	Pro	GIU	240	
173	223					230					233					240	
174	CCC	AAC	ACC	ACC	TCC	CCG	GAG	тст	CCC	GAC	ACC	ACC	TCC	CCG	GAG	тст	768
175							Glu										
176					245					250					255		
177																	
178							GAG										816
179	Pro	Asp	Thr		Ser	Gln	Glu	Pro		Asp	Thr	Thr	Ser		Glu	Pro	
180				260					265					270			
181 182	ccc	CAC	A C C	N.C.C	TCC	CAC	GAG	CCT	ccc	CAC	N.C.C	N.C.C	TOO	ccc	CAC	CCT	864
183							Glu										004
184	110	пор	275	****	DCI	0111	014	280	110	пор			285	110	Olu	110	
185																	
186	CCC	GAC	AAG	ACC	TCC	CCG	GAG	CCC	GCC	CCC	CAG	CAG	GGC	TCC	ACA	CAC	912
187	Pro	Asp	Lys	Thr	Ser	Pro	Glu	Pro	Ala	Pro	Gln	Gln	Gly	Ser	Thr	His	
188		290					295					300					
189																	
190							TCC										960
191 192	305	PIO	Arg	ser	PIO	310	Ser	THE	Arg	THE	315	Arg	Pro	GIU	тте	320	
193	303					310					313					320	
194	CAG	GCT	GGG	ccc	ACG	CAG	GGA	GAA	GTG	ATC	CCA	ACA	GGC	TCG	TCC	AAA	1008
195							Gly										
196			-		325		•			330			•		335	-	
197																	
198							CCC										1056
199	Pro	Ala	Gly	_	Gln	Leu	Pro	Ala		Leu	Trp	Thr	Ser		Ala	Val	
200				340					345					350			
201	OTT C	CCA	аша	OMO.	OTIC!	ama	aaa	mma	aaa	N.C.C	יחאות	C N C	оша	maa	***	aaa	1104
202 203							GCC Ala										1104
203	Leu	GTA	355	Leu	rea	red	MIG	360	FIO	1111	ı y i	птэ	365	тър	гур	ALG	
205			555														

258

RAW SEQUENCE LISTING PATENT APPLICATION US/08/875,849

DATE: 09/15/98 TIME: 15:05:59

	INDIA CIA	
006	INPUT SET: S2864	
206	TGC CGG CAC CTG GCT GAG GAC GAC ACC CAC CCA CCA GCT TCT CTG AGG	1152
207	Cys Arg His Leu Ala Glu Asp Asp Thr His Pro Pro Ala Ser Leu Arg	
208	370 375 380	
209		
210	CTT CTG CCC CAG GTG TCG GCC TGG GCT GGG TTA AGG GGG ACC GGC CAG	1200
211	Leu Leu Pro Gln Val Ser Ala Trp Ala Gly Leu Arg Gly Thr Gly Gln	
212	385 390 395 400	
213		
214	GTC GGG ATC AGC CCC TCC TGAGTGGCCA GCCTTTCCCC CTGTGAAAGC	1248
215	Val Gly Ile Ser Pro Ser	
216	405	
217		
218	AAAATAGCTT GGACCCCTTC AAGTTGAGAA CTGGTCAGGG CAAACCTGCC TCCCATTCTA	1308
219		
220	CTCAAAGTCA TCCCTCTGCT CACAGAGATG GATGCATGTT CTGATTGCCT CTTTGGAGAA	1368
221		
222	GCTCATCAGA AACTCAAAAG AAGGCCACTG TTTGTCTCAC CTACCCATGA CCTGAAGCCC	1428
223	detertered aneternano anocenero l'ilidiciene elicectator ceronacce	1420
224	CTCCCTGAGT GGTCCCCACC TTTCTGGACG GAACCACGTA CTTTTTACAT ACATTGATTC	1488
225	CICCUIGNOT GOICCCACC THICIGONG GAACCACGTA CITTITACAT ACATTGATIC	1400
225	ATGTCTCACG TCTCCCTAAA AATGCGTAAG ACCAAGCTGT GCCCTGACCA CCCTGGGCCC	1548
	AIGICICACE ICICCCIAAA AAIGCGIAAG ACCAAGCIGI GCCCIGACCA CCCIGGGCCC	1340
227	CHARGE CHARGE CACAMAGNA CACAMANAGA AND LAGRA CHARLANDA AND LAGRA CHARLANDA AND LAGRA CHARLANDA AND LAGRA CHARLANDA CHARGA CHARLANDA CHARCANDA CHARCA CHARCANDA CHARCAND	1.600
228	CTGTCGTCAG GACCTCCTGA GGCTTTGGCA AATAAACCTC CTAAAATGAT AAAAAAAAA	1608
229		
230	AAAAAAAA AAAAA	1624
	ABARBARBAR ABARBA	
231	ADDROGODO ADDROG	
231 232	pagagagaga angga	
231 232 233		
231 232 233 234	(2) INFORMATION FOR SEQ ID NO:2:	
231 232 233 234 235	(2) INFORMATION FOR SEQ ID NO:2:	
231 232 233 234 235 236	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS:	
231 232 233 234 235	(2) INFORMATION FOR SEQ ID NO:2:	
231 232 233 234 235 236 237 238	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid	
231 232 233 234 235 236 237 238 239	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids	
231 232 233 234 235 236 237 238	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid	
231 232 233 234 235 236 237 238 239	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid	
231 232 233 234 235 236 237 238 239 240	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear	
231 232 233 234 235 236 237 238 239 240 241	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear	
231 232 233 234 235 236 237 238 239 240 241 242	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear	
231 232 233 234 235 236 237 238 239 240 241 242 243	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear	
231 232 233 234 235 236 237 238 239 240 241 242 243 244	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: Met Asp Phe Gly Leu Ala Leu Leu Leu Ala Gly Leu Leu Gly Leu Leu	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: Met Asp Phe Gly Leu Ala Leu Leu Leu Ala Gly Leu Leu Gly Leu Leu 1 5 10 15	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: Met Asp Phe Gly Leu Ala Leu Leu Leu Ala Gly Leu Leu Gly Leu Leu 1 5 10 15 Leu Gly Gln Ser Leu Gln Val Lys Pro Leu Gln Val Glu Pro Pro Glu	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: Met Asp Phe Gly Leu Ala Leu Leu Leu Ala Gly Leu Leu Gly Leu Leu 1 5 10 15	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: Met Asp Phe Gly Leu Ala Leu Leu Leu Ala Gly Leu Leu Gly Leu Leu 1 5 10 15 Leu Gly Gln Ser Leu Gln Val Lys Pro Leu Gln Val Glu Pro Pro Glu 20 25 30	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: Met Asp Phe Gly Leu Ala Leu Leu Leu Ala Gly Leu Leu Gly Leu Leu leu lous lous lous lous lous lous lous lou	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253 254	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: Met Asp Phe Gly Leu Ala Leu Leu Leu Ala Gly Leu Leu Gly Leu Leu 1 5 10 15 Leu Gly Gln Ser Leu Gln Val Lys Pro Leu Gln Val Glu Pro Pro Glu 20 25 30	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 250 251 252 253 255	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: Met Asp Phe Gly Leu Ala Leu Leu Leu Ala Gly Leu Leu Gly Leu Leu 1 5 15 Leu Gly Gln Ser Leu Gln Val Lys Pro Leu Gln Val Glu Pro Pro Glu 20 25 30 Pro Val Val Ala Val Ala Leu Gly Ala Ser Arg Gln Leu Thr Cys Arg 40 45	
231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253 254	(2) INFORMATION FOR SEQ ID NO:2: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 406 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear (ii) MOLECULE TYPE: protein (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: Met Asp Phe Gly Leu Ala Leu Leu Leu Ala Gly Leu Leu Gly Leu Leu leu lous lous lous lous lous lous lous lou	

SEQUENCE VERIFICATION REPORT PATENT APPLICATION *US/08/875,849*

DATE: 09/15/98 TIME: 15:06:01

Line	Error	Original Text
5	Mandatory Value Not Present	(i) APPLICANT:
6	Unknown or Misplaced Identifier	(A) NAME: LeukoSite, Inc.
7	Unknown or Misplaced Identifier	(B) STREET: 215 First Street
8	Unknown or Misplaced Identifier	(C) CITY: Cambridge
9	Unknown or Misplaced Identifier	(D) STATE/PROVINCE: Massachusetts
10	Unknown or Misplaced Identifier	(E) COUNTRY: U.S.A.
11	Unknown or Misplaced Identifier	(F) POSTAL CODE/ZIP: 02142
12	Unknown or Misplaced Identifier	(G) TELEPHONE: (617) 621-9350
13	Unknown or Misplaced Identifier	(I) TELEFAX: (617) 621-9349
15	Mandatory Value Not Present	(i) APPLICANT/INVENTOR:
16	Unknown or Misplaced Identifier	(A) NAME: Michael J. Briskin
17	Unknown or Misplaced Identifier	(B) STREET: 28 Harbell Street
18	Unknown or Misplaced Identifier	(C) CITY: Lexington
19	Unknown or Misplaced Identifier	(D) STATE/PROVINCE: Massachusetts
20	Unknown or Misplaced Identifier	(E) COUNTRY: U.S.A.
21	Unknown or Misplaced Identifier	(F) POSTAL CODE/ZIP: 02173
23	Mandatory Value Not Present	(i) APPLICANT/INVENTOR:
24	Unknown or Misplaced Identifier	(A) NAME: Douglas J. Ringler
25	Unknown or Misplaced Identifier	(B) STREET: 382 Ocean Avenue, #1008
26	Unknown or Misplaced Identifier	(C) CITY: Revere
27	Unknown or Misplaced Identifier	(D) STATE/PROVINCE: Massachusetts
28	Unknown or Misplaced Identifier	(E) COUNTRY: U.S.A.
29	Unknown or Misplaced Identifier	(F) POSTAL CODE/ZIP: 02151
31	Mandatory Value Not Present	(i) APPLICANT/INVENTOR:
32	Unknown or Misplaced Identifier	(A) NAME: Dominic Picarella
33	Unknown or Misplaced Identifier	(B) STREET: 2 North Bennet Court, #4
34	Unknown or Misplaced Identifier	(C) CITY: Boston
35	Unknown or Misplaced Identifier	(D) STATE/PROVINCE: Massachusetts
36	Unknown or Misplaced Identifier	(E) COUNTRY: U.S.A.
37	Unknown or Misplaced Identifier	(F) POSTAL CODE/ZIP: 02113
39	Mandatory Value Not Present	(i) APPLICANT/INVENTOR:
40	Unknown or Misplaced Identifier	(A) NAME: Walter Newman
41	Unknown or Misplaced Identifier	(B) STREET: 3 Durham Street, #3
42	Unknown or Misplaced Identifier	(C) CITY: Boston
43	Unknown or Misplaced Identifier	(D) STATE/PROVINCE: Massachusetts
44	Unknown or Misplaced Identifier	(E) COUNTRY: U.S.A.
45	Unknown or Misplaced Identifier	(F) POSTAL CODE/ZIP: 02115